S3 Table. Univariable logistic regression models for *Toxoplasma gondii* seropositivity for cats of eight breeds separately as well as for all cats, regardless of breed. Odds ratio, 95% confidence interval, and P-value are shown.

-7.07), P=0.066 1.10 -2.78), P=0.573 1.88	3 (0.33–0.87), P=0.012 0 (0.48–2.52), P=0.819 8 (0.63–5.62), P=0.262 9 (0.35-1.78), P=0.572	1.42 (0.35–5. 8.75 (1.08–70	71), P=0.622 0.70), P=0.042	1.23 (0.75–2.02), P=0.416 0.80 (0.34–1.85), P=0.597 1 (0.32–3.15), P=1.000
-2.78), P=0.573 1.88	8 (0.63–5.62), P=0.262	8.75 (1.08–70		, , , , , , , , , , , , , , , , , , ,
,	. , , , , , , , , , , , , , , , , , , ,	,	0.70), P=0.042	1 (0.32–3.15), P=1.000
-9.49), P=0.415 0.79	9 (0.35-1.78), P=0.572			
	, (3.20 2.7.0), 2 3.072	1.78 (0.77-4.3	11), P=0.178	0.71 (0.28-1.78), P=0.460
-8.53), P=0.000 0.93	5 (0.62–1.45), P=0.809	1.39 (0.63–3.	.07), P=0.409	2.06 (1.20–3.53), P=0.008
) 1.6	1 (0.69–3.78), P=0.270	0.78 (0.05–12	2.95), P=0.864	1.65 (0.69–3.91), P=0.259
0.84	4 (0.30–2.39), P=0.750	1.38 (0.18–10	0.61), P=0.756	1.82 (0.58–5.71), P=0.306
0.65	5 (0.16–2.60), P=0.548	1 (omitted)		0.25 (0.06–1.08), P=0.064
-5.38), P=0.000 0.83	8 (0.69–1.12), P=0.290	2.58 (1.78–3.	.74), P=0.000	1.43 (1.11–1.83), P=0.005
) 1.6) 0.8) 0.6	1.61 (0.69–3.78), P=0.270 0.84 (0.30–2.39), P=0.750 0.65 (0.16–2.60), P=0.548	1.61 (0.69–3.78), P=0.270	1.61 (0.69–3.78), P=0.270